1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/530,746A

DATE: 08/28/2001 TIME: 08:01:02

Input Set : A:\1803-277-999.txt

Output Set: N:\CRF3\08282001\1530746A.raw

	sacpae see/exi3/ssssssssssssssssssssssss	ENTERED
4	<110> APPLICANT: Kessler, Christoph	ENIERED
5		
6	Bartl, Knut	
7	Orum, Henrik	
9	<120> TITLE OF INVENTION: SPECIFIC AND SENSITIVE METHOD	FOR DETECTING NUCLEIC ACIDS
12	<130> FILE REFERENCE: 4817/OQ	
C> 14	<140> CURRENT APPLICATION NUMBER: US/09/530,746A	
C> 15	<141> CURRENT FILING DATE: 2000-11-16	
17	<160> NUMBER OF SEQ ID NOS: 94	
19	<170> SOFTWARE: PatentIn Version 3.1	
21	<210> SEQ ID NO: 1	
22	<211> LENGTH: 24	
23	<212> TYPE: DNA	
24	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
27	<223> OTHER INFORMATION: amplification primer	
29	<400> SEQUENCE: 1	
30	gcagaaagcg tctagccatg gcgt	24
33	<210> SEQ ID NO: 2	
34	<211> LENGTH: 24	
35	<212> TYPE: DNA	
36	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
39	<223> OTHER INFORMATION: amplification primer	
41	<400> SEQUENCE: 2	
42	ctcgcaagca ccctatcagg cagt	24
45	<210> SEQ ID NO: 3	
46	<211> LENGTH: 20	
47	<212> TYPE: DNA	
48	<213> ORGANISM: Artificial Sequence	
50	<220> FEATURE:	
51	<223> OTHER INFORMATION: amplification primer	
53	<400> SEQUENCE: 3	
54	agtatgtgtg tcgtgcagcc	20
57	<210> SEQ ID NO: 4	
58	<211> LENGTH: 18	
59	<212> TYPE: DNA	
60	<213> ORGANISM: Artificial Sequence	
62	<220> FEATURE:	
63	<223> OTHER INFORMATION: amplification primer	
	<400> SEQUENCE: 4	
	tggctctccc gggagtgg	18
	<210> SEQ ID NO: 5	
	<211> LENGTH: 12	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
74	<220> FEATURE:	

DATE: 08/28/2001

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/530,746A TIME: 08:01:02

Input Set : A:\1803-277-999.txt

Output Set: N:\CRF3\08282001\I530746A.raw	
75 <223> OTHER INFORMATION: probe	
77 <400 > SEQUENCE: 5	
78 otomaggaco oc	12
81 <210> SEQ ID NO: 6	
82 <211> LENGTH: 48	
83 <212> TYPE: DNA	
84 (213> ORGANISM: HCV	
86 - 400> SEQUENCE: 6	
87 adtatgagtg tegtgeagee teeaggaeee eeeeteeegg gagageea	48
90 -210> SEQ ID NO: 7	
91 -211> LENGTH: 48	
92 <212> TYPE: DNA	
93 (213> ORGANISM: Human	
95 <400> SEQUENCE: 7	
96 agtatgtgtg togtgoagoo tocaggacoo coactooogg gagagooa	48
99 <210> SEQ ID NO: 8	
100 <211> LENGTH: 59	
101 <212> TYPE: DNA	
102 <213> ORGANISM: HCV	
104 <400> SEQUENCE: 8	
105 gtactgcctg atagggtgct tgcgagtgcc ccgggaggtc tcgtagaccg tgcaccatg	59
108 <210> SEQ ID NO: 9	
109 <211> LENGTH: 59	
110 <212> TYPE: DNA	
111 <213> ORGANISM: HGBV-B	
113 <400> SEQUENCE: 9	
114 gtactgcctg atagggtcct tgcgagggga tctgggagtc tcgtagaccg tagcacatg	59
117 <210> SEQ ID NO: 10	
118 <211> LENGTH: 19	
119 <212> TYPE: DNA	

- 122 <220> FEATURE: 123 <223> OTHER INFORMATION: amplification primer
- 125 <400> SEQUENCE: 10
- 126 ccaggacccc cactcccgg 19
- 129 <210> SEQ ID NO: 11
- 130 <211> LENGTH: 20
- 131 <212> TYPE: DNA
- 132 <213> ORGANISM: Artificial Sequence

120 <213> ORGANISM: Artificial Sequence

- 134 <220> FEATURE:
- 135 <223> OTHER INFORMATION: amplification primer
- 137 <400> SEQUENCE: 11
- 138 tocaggacco coactoogg
- 141 <210> SEQ ID NO: 12
- 142 <211> LENGTH: 16
- 143 <212> TYPE: DNA
- 144 <213> ORGANISM: Artificial Sequence
- 146 <220> FEATURE:
- 147 <223> OTHER INFORMATION: amplification primer

20

RAW SEQUENCE LISTING

DATE: 08/28/2001

PATENT APPLICATION: US/09/530,746A

TIME: 08:01:02

Input Set : A:\1803-277-999.txt

Output Set: N:\CRF3\08282001\1530746A.raw

	<.400> SEQUENCE: 12	
	deaggaeded cacted	16
	-210> SEQ ID NO: 13	
	C2115 LENGTH: 48	
	<pre><212> TYPE: DNA </pre>	
	+2213> ORGANISM Artificial Sequence	
	<pre><220> FEATURE: <223> OTHER INFORMATION: amplification primer</pre>	
	<2250 OTHER INFORMATION: dispitited closs primer <400> SEQUENCE: 13	
	autatgagtg tegtgeagee tecaggeece eccetecegg gagageea	48
	<210> SEQ ID NO: 14	
	<211> LENGTH: 22	
	<212> TYPE: DNA	
	<pre><213> ORGANISM: Artificial Sequence</pre>	
	<220> FEATURE:	
171	<223> OTHER INFORMATION: amplification primer	
173	<pre><400> SEQUENCE: 14</pre>	
	gtgtgtcgtg cagcctccag ga	22
177	<210> SEQ ID NO: 15	
	<211> LENGTH: 17	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<pre>< 220> FEATURE:</pre>	
	< 223> OTHER INFORMATION: amplification primer	
	.400> SEQUENCE: 15	17
	togtgcagoo tocagga <210> SEQ ID NO: 16	1 /
	<211> LENGTH: 18	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION: amplification primer	
	<400> SEQUENCE: 16	
198	ccactcccgg gagageca	18
201	<210> SEQ ID NO: 17	
	<211> LENGTH: 21	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<pre><223> OTHER INFORMATION: amplification primer</pre>	
	<400> SEQUENCE: 17	2.1
	cqtactgcct gatagggtgc t	21
	<210> SEQ ID NO: 18 <211> LENGTH: 23	
	<pre><211> LENGTH: 23 <212> TYPE: DNA</pre>	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	

221 <400> SEQUENCE: 18

219 <223> OTHER INFORMATION: amplification primer

RAW SEQUENCE LISTING

DATE: 08/28/2001 TIME: 08:01:02

PATENT APPLICATION: US/09/530,746A

Input Set : A:\1803-277-999.txt

Output Set: N:\CRF3\08282001\1530746A.raw

			tgcta mggt.mtamga gac	23
			SEQ ID NO: 19	
			LENGTH: 21	
			TYPE: DNA	
			ORGANISM: Artificial Sequence	
			FEATURE:	
			OTHER INFORMATION: amplification primer	
			SEQUENCE: 19	0.3
			tgoot gatagggttg c	21
			SEQ ID NO: 20	
			LENGTH: 23	
			TYPE: DNA	
			ORGANISM: Artificial Sequence	
			FEATURE:	
			OTHER INFORMATION: amplification primer	
			SEQUENCE: 20	0.3
			tgmta mggtmtamga gac	23
			SEQ ID NO: 21	
			LENGTH: 21	
			TYPE: DNA	
			ORGANISM: Artificial Sequence	
			FEATURE:	
			OTHER INFORMATION: amplification primer	
r.r \ \	207	<222U>	FEATURE:	
W-FS	258	4221 >	NAME/KEY: modified base LOCATION: 11,19	
			OTHER INFORMATION: n=i SEQUENCE: 21	
Tat u N			tgcct natagggtnc t	21
12E			SEQ ID NO: 22	21
			LENGTH: 23	
			TYPE: DNA	
			ORGANISM: Artificial Sequence	
			FEATURE:	
			OTHER INFORMATION: amplification primer	
			FEATURE:	
4-L			NAME/KEY: modified base	
.			LOCATION: 19	
			OTHER INFORMATION: n=i	
			SEQUENCE: 22	
w/~>	280	amatat	tgmta mggtmtdmnd gdc	23
.	283	<210>	SEQ ID NO: 23	
			LENGTH: 20	
			TYPE: DNA	
			ORGANISM: Artificial Sequence	
			FEATURE:	
			OTHER INFORMATION: amplification primer	
			FEATURE:	
W>			NAME/KEY: modified base	
••		,		

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/530,746A II

DATE: 08/28/2001 TIME: 08:01:02

Input Set : A:\1803-277-999.txt

Output Set: N:\CRF3\08282001\I530746A.raw

293 <222> LOCATION: 11,19 294 -223> OTHER INFORMATION: n=i 296 <400> SEQUENCE: 23 20 v√-> 297 cgtactgcct natagggtnc 300 <210> SEQ ID NO: 24 301 <211> LENGTH: 21 302 - 1212> TYPE: DNA 303 <213> ORGANISM: Artificial Sequence 305 -220> FEATURE: 306 - 223> OTHER INFORMATION: amplification primer 308 +.220> FEATURE: W > 309 <221> NAME/KEY: modified base 310 <222> LOCATION: 11,19 311 <223> OTHER INFORMATION: n=i 313 <400> SEQUENCE: 24 21 w-> 314 cgtamtgmmt natagggtnm t 317 <210> SEQ ID NO: 25 318 <211> LENGTH: 23 319 (212> TYPE: DNA 320 <213> ORGANISM: Artificial Sequence 322 <220> FEATURE: 323 <223> OTHER INFORMATION: amplification primer 325 <220> FEATURE: ₩--> 326 <221> NAME/KEY: modified base 327 <222> LOCATION: 19 328 <223> OTHER INFORMATION: n=i 330 <220> FEATURE: ₩₩> 331 <221> NAME/KEY: modified base 332 <222> LOCATION: 6,8,9 333 <223> OTHER INFORMATION: n=p335 <400> SEQUENCE: 25 wi->⇒ 336 gmatgnknna mggtmtdmnd gdm 23 339 <210> SEQ ID NO: 26 340 <211> LENGTH: 20 341 <212> TYPE: DNA 342 <213> ORGANISM: Artificial Sequence 344 <220> FEATURE: 345 <223> OTHER INFORMATION: amplification primer 347 <220> FEATURE: (W-> 348 <221> NAME/KEY: modified base 349 <222> LOCATION: 11,19 350 <223> OTHER INFORMATION: n=i 353 <400> SEQUENCE: 26 ₩\→> 354 cgtamtgmmt natagggtnm 20 357 <210> SEQ ID NO: 27 358 <211> LENGTH: 26 359 <212> TYPE: DNA 360 <213> ORGANISM: Artificial Sequence

The second of th

362 <220> FEATURE:

VERIFICATION SUMMARY

DATE: 08/28/2001 PATENT APPLICATION: US/09/530,746A TIME: 08:01:04

Input Set : A:\1803-277-999.txt

Output Set: N:\CRF3\08282001\I530746A.raw

L.15 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:258 M:257 W: Feature value mis-spelled or invalid, <2215 Name/Key for SEQ ID#:21 L:263 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 L:275 M:257 W: Feature value mis-spelled or invalid, <2215 Name/Key for SEQ ID#:22 L:280 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 L:292 M:257 W: Feature value mis-spelled or invalid, <2215 Name/Key for SEQ ID#:23 L:297 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 L:309 M:257 W: Feature value mis-spelled or invalid, <2215 Name/Key for SEQ ID#:24 L:314 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 L:326 M:257 W: Feature value mis-spelled or invalid, <2215 Name/Key for SEQ ID#:25
L:280 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 L:292 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:23 L:297 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 L:309 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:24 L:314 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L.297 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 L:309 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:24 L:314 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L 314 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L.326 M:257 W: Feature value mis-spelled or invalid, +221> Name/Key for SEQ 1D#:25
L:331 M.257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:25
L:336 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:348 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:26 L:354 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:378 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:28 L:384 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:396 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:29 L:402 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:415 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:30 L:420 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:432 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:31
L:437 M:257 W: Feature value mis-spelled or invalid, $<221>$ Name/Key for SEQ ID#:31 L:443 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:455 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:32
L:461 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:32
L:466 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:502 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:35 L:507 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:519 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:36
L:524 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36
L:536 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:37
L:541 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:37
L:547 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37
L:559 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:38
L:564 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:38
L:569 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38
L:581 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:39
L:586 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39
L:599 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:40
L:604 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:40
L:609 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40
L:621 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:41
L:626 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:41 L:631 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41
L:655 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:43
L:660 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43
L:672 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:44

VERIFICATION SUMMARY

L.770 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49

PATENT APPLICATION: US/09/530,746A TIME: 08:01:04

DATE: 08/28/2001

Input Set : A:\1803-277-999.txt

Output Set: N:\CRF3\08282001\1530746A.raw

L:677 M:341 W:	(46) "n" or "Xaa" used, for SEQ ID#:44				
L:691 M:257 W:	Feature value mis-spelled or invalid, <221>	Name/Key	for	SEQ	ID#:45
L:696 M:257 W:	Feature value mis-spelled or invalid, <221>	Name/Key	for	SEQ	ID#:45
L:702 M:341 W:	(46) "n" or "Xaa" used, for SEQ ID#:45				
L:726 M:257 W:	Feature value mis-spelled or invalid, <2215	Name/Key	for	SEQ	ID#:47
L.731 M:341 W:	(46) "n" or "Xaa" used, for SEQ ID#:47				
L:743 M:257 W:	Feature value mis-spelled or invalid, <221>	Name/Key	for	SEQ	ID#:48
L.748 M:341 W:	(46) "n" or "Xaa" used, for SEQ ID#:48				
L:760 M:257 W:	Feature value mis-spelled or invalid, <221>	Name/Key	for	SEQ	ID#:49
L:765 M:257 W:	Feature value mis-spelled or invalid, <221>	Name/Key	for	SEQ	ID#:49